SCORE Search Results Details for Application 08900220 and Search Result us-08-900-220c-17.rai.

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SCORE System Overview

SCORE FAQ Comments / Suggestions

This page gives you Search Results detail for the Application 08900220 and Search Result us-08-900-220c-17.rai.

start

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OM protein - protein search, using sw model

Run on:

May 25, 2006, 11:40:16 ; Search time 50 Seconds

(without alignments)
693.243 Million cell updates/sec

Title:

US-08-900-220C-17

Perfect score: 2088

Sequence: 1

1 MALLTNLLPLCCLALLALPA......pTGMHWYSRLLYRLAEELLG 396

Scoring table:

BLOSUM62

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Searched:

650591 seqs, 87530628 residues

Total number of hits satisfying chosen parameters:

650591

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

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Pred. No. is the number of results predicted by chance to have a

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ALIGNMENTS

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RESULT 1
US-09-325-256-24
; Sequence 24, Application US/09325256
 Patent No. 6444793
 GENERAL INFORMATION:
  APPLICANT: PEPINSKY, R. BLAKE
  APPLICANT: BAKER, DARREN P.
  APPLICANT: WEN, DINGYI
  APPLICANT: WILLIAMS, KEVIN P.
  APPLICANT: GARGER, ELLEN A.
  APPLICANT: TAYLOR, FREDERICK R.
  APPLICANT: GALDES, ALPHONSE
  APPLICANT: PORTER, JEFFREY
  TITLE OF INVENTION: HYDROPHOBICALLY-MODIFIED PROTEIN COMPOSITIONS AND
  TITLE OF INVENTION:
                       METHODS
  FILE REFERENCE: BIV-067.01
  CURRENT APPLICATION NUMBER: US/09/325,256
  CURRENT FILING DATE: 1999-06-03
  PRIOR APPLICATION NUMBER: 60/099,800
  PRIOR FILING DATE: 1998-09-10
  PRIOR APPLICATION NUMBER: 60/078,935
  PRIOR FILING DATE:
                      1998-03-20
  PRIOR APPLICATION NUMBER:
  PRIOR FILING DATE: 1998-06-17
  PRIOR APPLICATION NUMBER: 60/067,423
  PRIOR FILING DATE: 1997-12-03
  PRIOR APPLICATION NUMBER: PCT/US98/25676
  PRIOR FILING DATE:
                      1998-12-03
  NUMBER OF SEQ ID NOS: 31
  SOFTWARE: PatentIn Ver. 2.1
 SEQ ID NO 24
   LENGTH: 396
   TYPE: PRT
   ORGANISM: Homo sapiens
US-09-325-256-24
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OM protein - protein search, using sw model

Run on: May 25, 2006, 11:51:45; Search time 181 Seconds (without alignments)

1013.442 Million cell updates/sec

Title: US-08-900-220C-17

Perfect score: 2088

Sequence: 1 MALLTNLLPLCCLALLALPA......PTGMHWYSRLLYRLAEELLG 396

Scoring table: BLOSUM62

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Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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27	2008	96.2	396	3	US-09-021-660A-35	Sequence 35, Appl
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OM protein - protein search, using sw model

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(without alignments)

294.026 Million cell updates/sec

Title: US-08-900-220C-17

Perfect score: 2088

Sequence: 1 MALLTNLLPLCCLALLALPA.....PTGMHWYSRLLYRLAEELLG 396

Scoring table: BLOSUM62

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